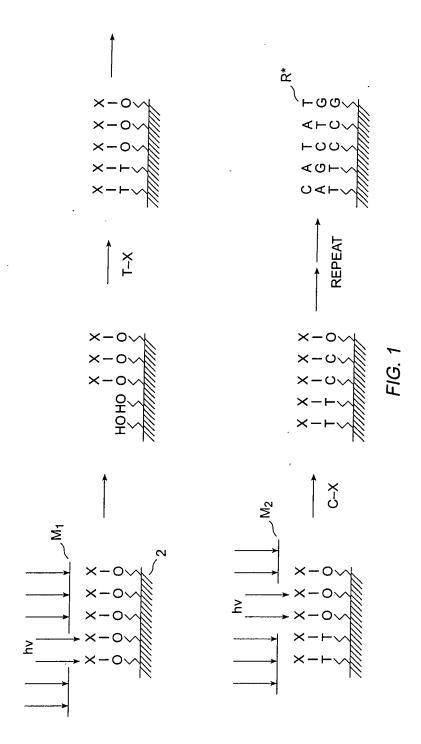
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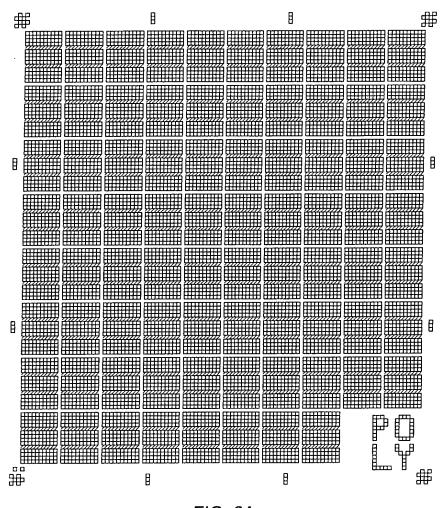


FIG. 2A

HOTHER OFFORGED

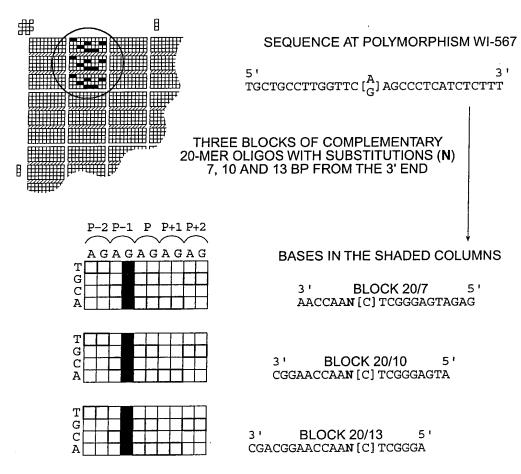
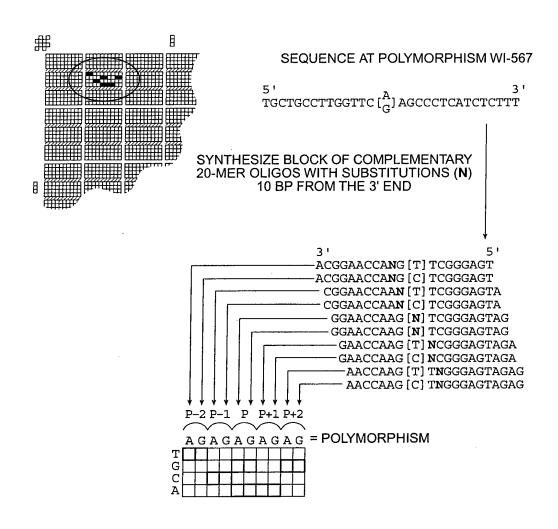


FIG. 2B



## PREDICTED PATTERNS

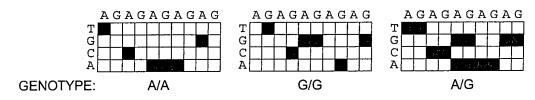


FIG. 3

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APPROVED O.G. FIG.

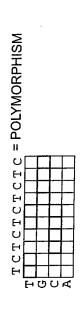
BY CLASS SUBCLASS

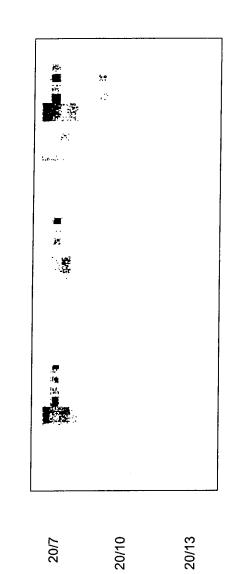
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 $WI-1959 = \dots ACCAAAAATCAGTC[T/C]GGGTAACTGAGAGTG...$ 





H B D A	
T C T C T C T C T C C C C C C C C C C C	C/C HOMOZYGOTE
TCTCTCTCTCGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	T/C HETEROZYGOTE

FIG 4

T/T HOMOZYGOTE

TCTCTCTCT

HOTHO, CHICOCO

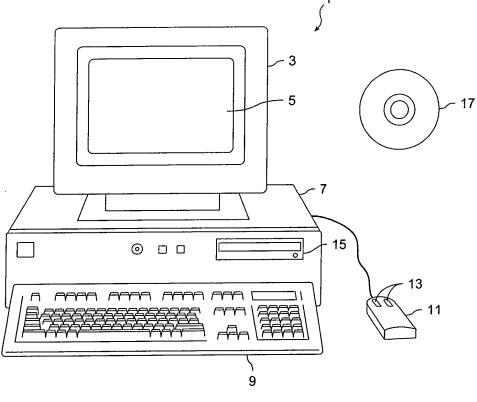


FIG. 5

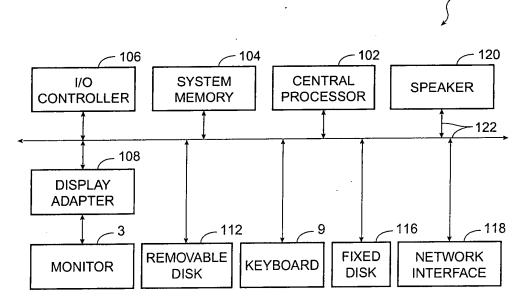


FIG. 6

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APPROVED O.G. FIG.

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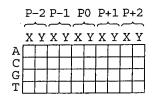


FIG. 7

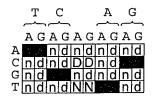


FIG. 8

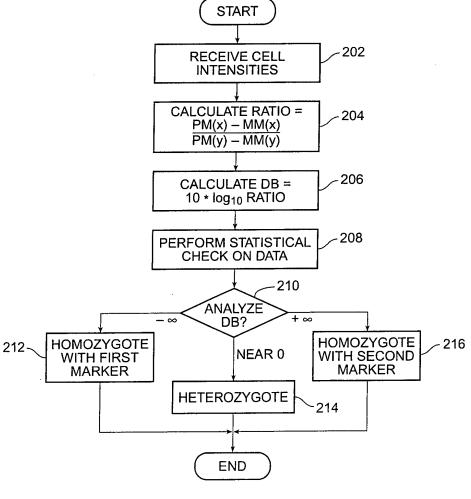
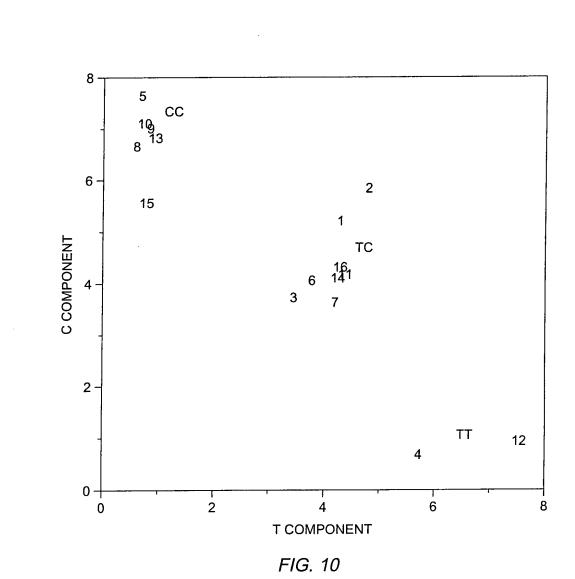


FIG. 9

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APPHOVED O.G. FIG.
BY CLASS SUBC

DOSSIL CEECT

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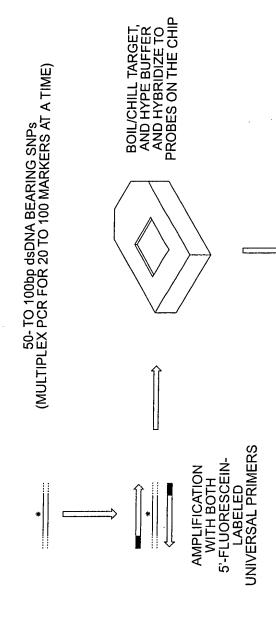


FIG. 11

SCAN AND ANYLYZE SIGNAL INTENSITIES

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DB = "DECIBELS"

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= 10 X LOG (DETECTION RATIO)

**DETECTION RATIO** 

A PERFECT MATCH - A MISMATCHES

B PERFECT MATCH - B MISMATCHES

C =

SIGNIFICANCE CHECK = MAX (A OR B PERFECT MATCH) > 2 σ (A OR B MISMATCHES)

PATTERN MATCHING

= MEAN [MAX (A OR B PERFECT MATCHES)] > 1.2 MEAN [MAX(A OR B MISMATCHES)]

PREDICTED PATTERNS

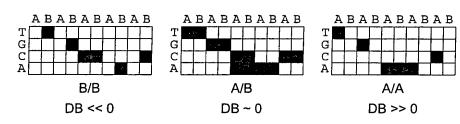


FIG. 12

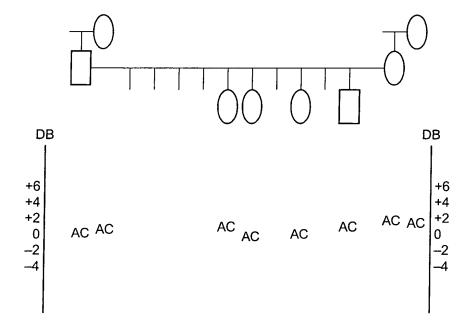


FIG. 13